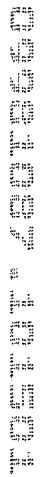
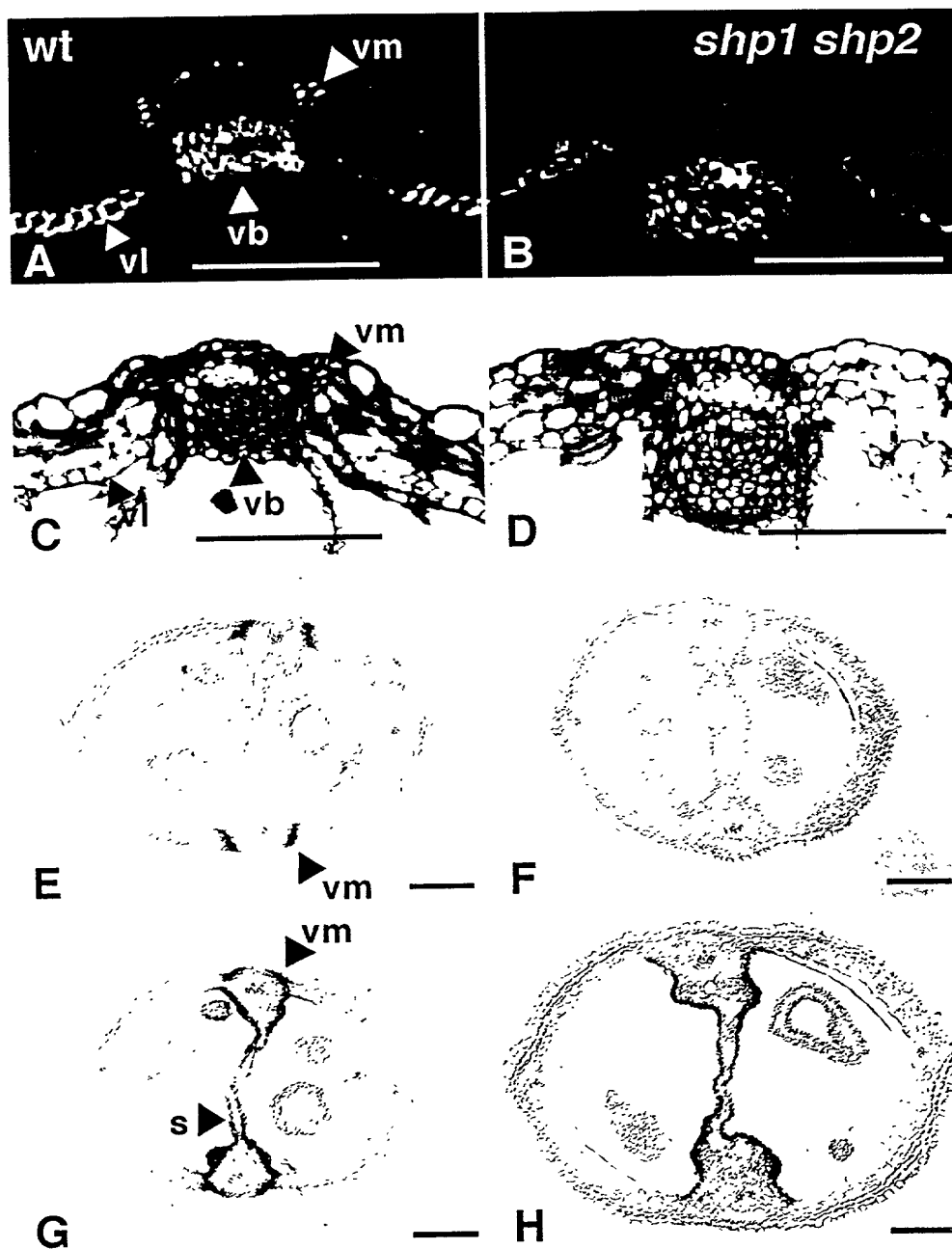
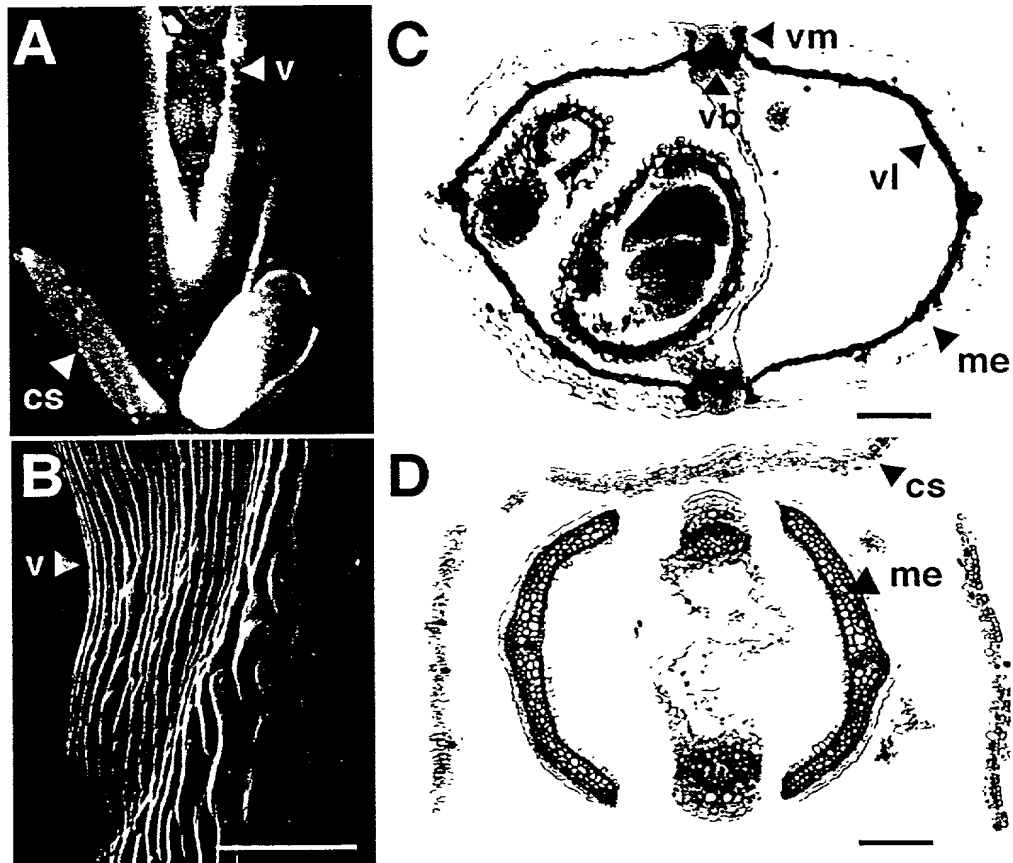
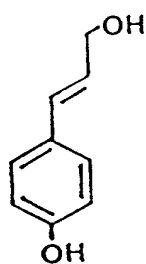


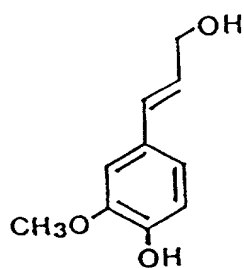
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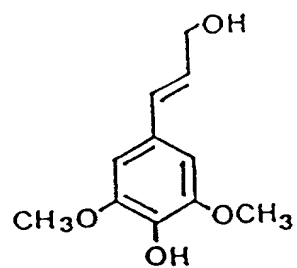




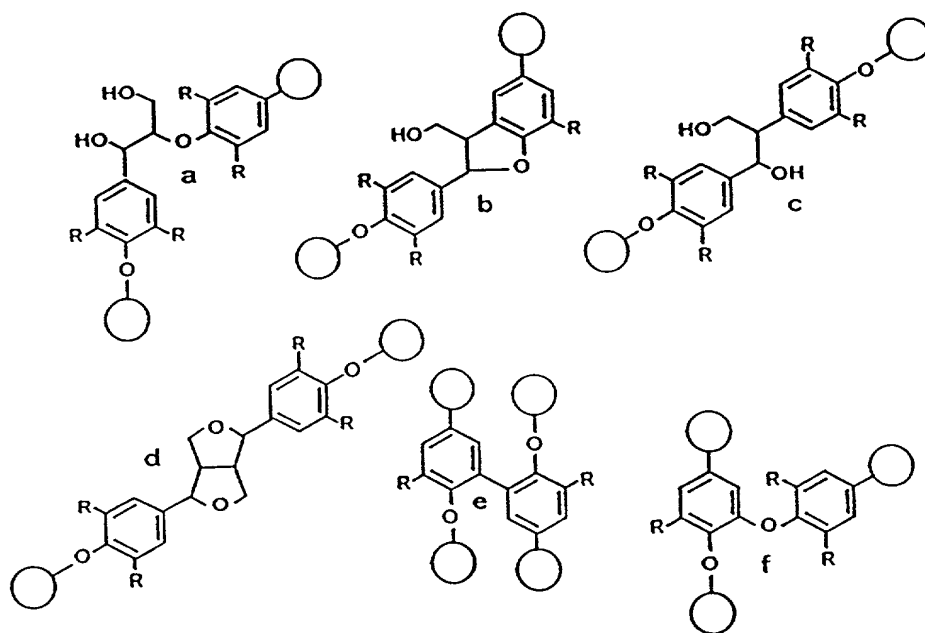
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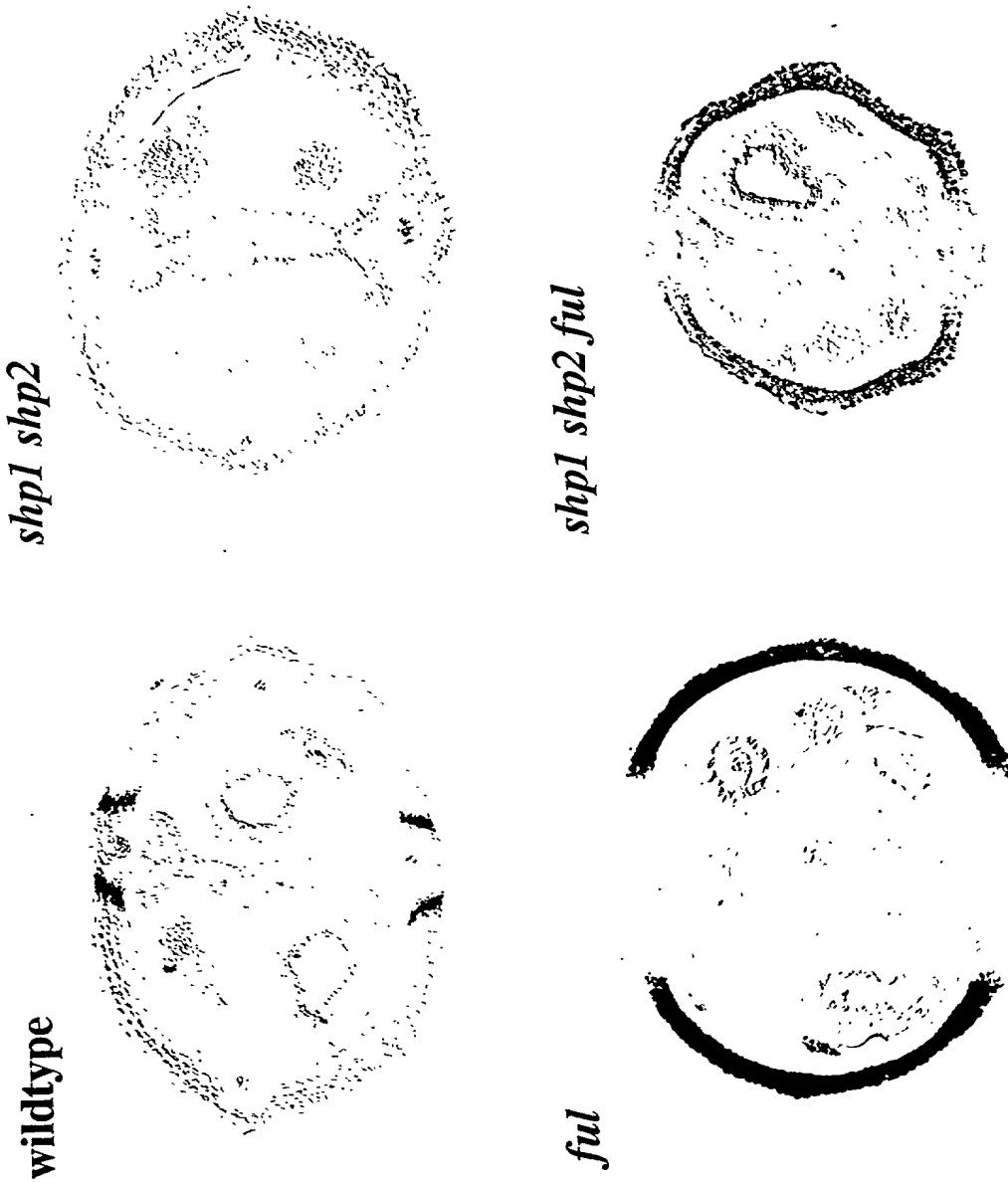


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3





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1 M G R G R V O L K R I E N K I N R O V T

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121 GCTGAGGTTCCTCTCATGCTCTTCTCTTCAAGGCAAACTCTTGAATATTCCACCGAC
41 A E V A L I V F S S K G K L F E Y S T D

181 TCTTCATGAGAGAGGATCTTGAAGCTATGATGCTATTTATATTTCAGACAAACAATT
61 S C H E R I L E R Y D R Y L Y S D K Q L

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81 V G R D V S Q S E N W Y L E H A K L K A

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3558-4-20-77

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